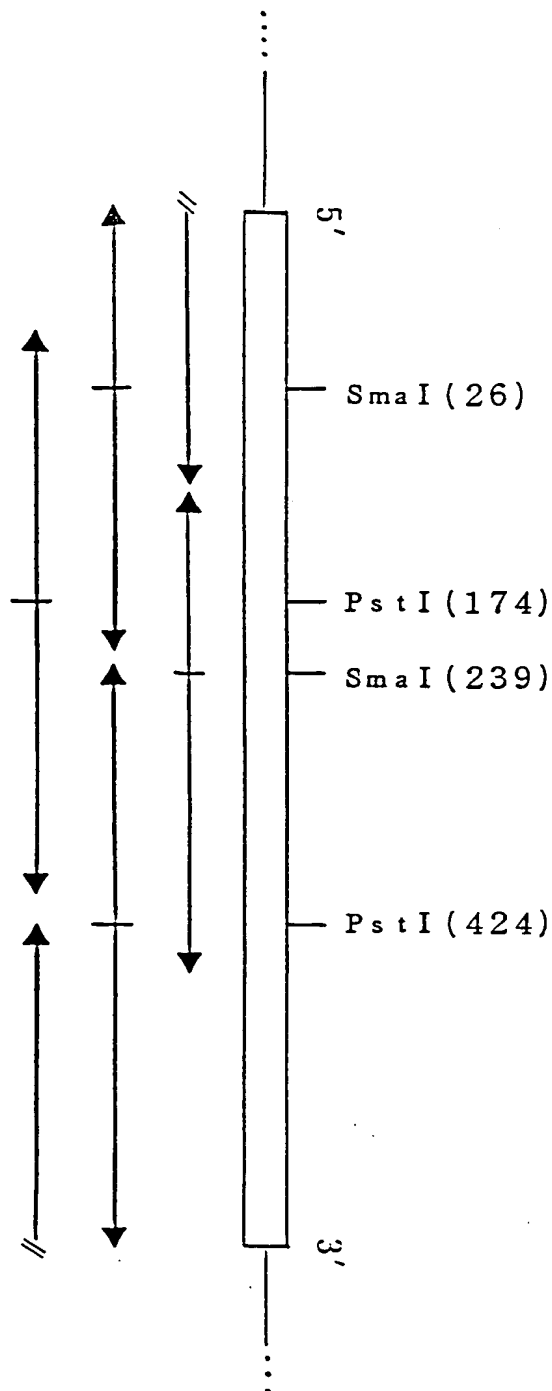


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FIG. 1



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99 GGCAAGGCTCAGGGCGAAGGTGGGAAGCAAACCCGGAGGCATCGCAGCACGCCAGCAGAACAGCAGCAACAGCCCTCCGCCATCTCCCTCCAGAGAC -1

90
ATG GAT CCC CAG ACA CCA CCT TCC CCG GCG CTC CTC CTC TTT TTT CAT CTG GCT TTC CTG GCA GAT GAT TCC AAC CCG CTG GGC
Het Asp Pro Gln Thr Ala Pro Ser Arg Ala Leu Leu Leu Leu Phe Leu His Leu Ala Phe Leu Gly Gly Arg Ser His Pro Leu Gly

180

270
CAC ACA TCC CTG CAG CCC CTC CAG GAG AGC CCC CGT CCC ACA GGT GTC TGG AAG TCC CGG GAG GTA GCC ACC GAG GGC ATC CGT GGC CAC
Gln Thr Ser Leu Glu Pro Leu Gln Glu Ser Pro Arg Pro Thr Gly Val Trp Lys Ser Arg Glu Val Ala Thr Glu Gly Ile Arg Gly His

CGC AAA ATG GTC CTC TAC ACC CTG CCG GCA CCA CGA AGC CCC AAG ATG GTG CAA GGG TCT GGC TCC TTT GGG AGG AAC ATG GAC CGG ATC
Arg Lys Met Val Leu Tyr Thr Leu Arg Ala Pro Arg Ser Pro Lys Met Val Gln Gly Ser Gly Arg Lys Met Asp Arg Ile

AGC TCC TCC AGT GGC CTG GGC TGC AAA GTG CTG AGG CCG CAT TAAAGAGGAAGTCTCTGGCTGCAGACACCTCTCTCTGATTTCCACAGGGGGCTTTTTCCTCAACCG
 Ser Ser Ser Ser Gly Leu Gly Cys Lys Val Leu Arg Arg His

584
TGTGGCGCCCTTTGAACTGACTCAATTTTTTAATGTAATTATGTAATTTTGAATGTTTATATAAGATGGTTTCTTACGTTTGAGCACAATTTCCACGGTCAAAATAAAGTCACACA

593 TTATTAAGCT

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FIG. 3-A

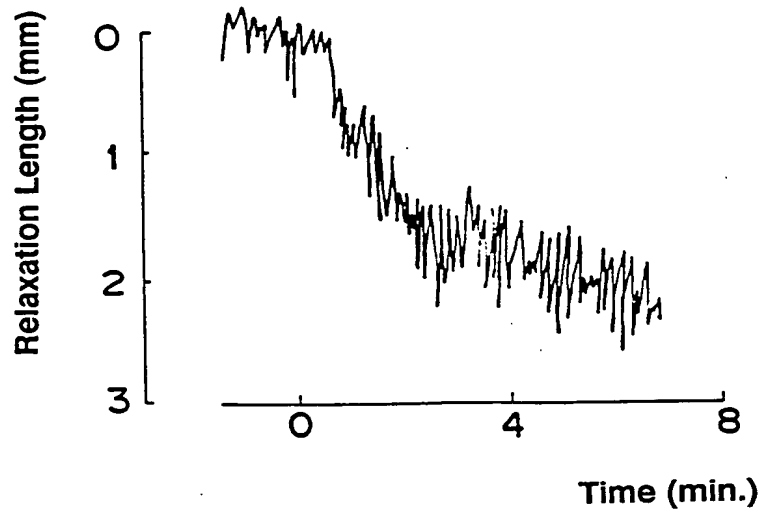


FIG. 3-B

